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MIRAGE (TM)  
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Release 3.1A John F. Collins, Biocomputing Research Unit.  
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Mpsrch\_nun n.a. - n.a. database search, using Smith-Waterman algorithm

Run on: Tue Oct 26 08:02:34 1999: Maspar time 23.19 Seconds  
940.454 Million cell updates/sec

Regular output not generated.

Description: >US-08-978-217-6

Perfect Score: (1-252) from US08978217.seq

N.A. Sequence: 252 1 AATTGTCCTTGAGAGCT.....CGGACACTGTGGCGACGA 252

Comp: TTAACAGCGGAGACTCTCGA.....GGCCGTGACACCGCGTCT

Scoring table: TABLE default

Gap 6

Nmatch STD: Dbase 0: Query 0

Searched: 165353 seqs, 43264542 bases x 2

Post-processing: Minimum Match 0%

Listing first 45 summaries

Database: n-Issued  
1:5A\_COMB 2:5B\_COMB 3:5C\_COMB 4:PCT9\_COMB 5:Backfiles1

Statistics: Mean 7.464: Variance 4.584: scale 1.628

Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

## SUMMARIES

o.	Score	Query Match	Length	DB	ID	Description	Pred. No.
1	252	100.0	1920	3	US-08-746-	Sequence 1, Applicatio	4.97e-152
2	34	13.5	215	1	US-08-238-	Sequence 5, Applicatio	1.77e-07
3	33	13.1	965	3	US-08-388-	Sequence 22, Applicati	6.53e-07
4	30	11.9	965	3	US-08-388-	Sequence 22, Applicati	3.06e-05
5	29	11.5	215	1	US-08-238-	Sequence 5, Applicatio	1.08e-04
6	26	10.3	74	4	PCT-US95-1	Sequence 94, Applicati	4.36e-03
7	26	10.3	81	4	PCT-US95-1	Sequence 92, Applicati	4.36e-03
8	24	9.5	75	4	PCT-US95-1	Sequence 99, Applicati	4.77e-02
9	24	9.5	82	4	PCT-US95-1	Sequence 97, Applicati	4.77e-02
10	24	9.5	2380	3	US-08-572-	Sequence 144, Applicat	1.54e-01
11	23	9.1	66	1	US-08-471-	Sequence 143, Applicat	1.54e-01
12	23	9.1	69	1	US-08-471-	Sequence 142, Applicat	1.54e-01
13	22	8.7	68	1	US-08-471-	Sequence 140, Applicat	4.85e-01
14	22	8.7	74	4	PCT-US95-1	Sequence 100, Applicat	4.85e-01
15	22	8.7	81	4	PCT-US95-1	Sequence 98, Applicati	4.85e-01
16	22	8.7	7559	3	US-08-250-	Sequence 2, Applicatio	4.85e-01
17	21	8.3	65	1	US-08-471-	Sequence 145, Applicat	1.50e+00
18	21	8.3	66	1	US-08-471-	Sequence 144, Applicat	1.50e+00
19	21	8.3	69	1	US-08-471-	Sequence 142, Applicat	1.50e+00
20	21	8.3	74	4	PCT-US95-1	Sequence 94, Applicati	1.50e+00

RESULT	ID	US-08-746-789A-1	STANDARD:	DNA:	UNC:	1920	BP.
21	21	8.3	81	4	PCT-US95-1	Sequence 92, Applicati	1.50e+00
22	21	8.3	82	4	PCT-US95-1	Sequence 97, Applicati	1.50e+00
23	21	8.3	1244	1	US-08-417-	Sequence 7, Applicatio	1.50e+00
24	21	8.3	1244	1	US-07-816-	Sequence 7, Applicatio	1.50e+00
25	21	8.3	2159	3	PCT-US95-1	Sequence 2, Applicatio	1.50e+00
26	21	8.3	2304	3	US-08-464-	Sequence 1, Applicatio	1.50e+00
27	21	8.3	2304	3	US-08-464-	Sequence 1, Applicatio	1.50e+00
28	21	8.3	6306	1	US-08-467-	Sequence 3, Applicatio	1.50e+00
29	21	8.3	6306	1	US-08-467-	Sequence 3, Applicatio	1.50e+00
30	21	8.3	6306	4	PCT-US93-0	Sequence 3, Applicatio	1.50e+00
31	21	8.3	6306	2	US-08-470-	Sequence 3, Applicatio	1.50e+00
32	21	8.3	6306	2	US-08-195-	Sequence 3, Applicatio	1.50e+00
33	21	8.3	6306	1	PCT-US95-1	Sequence 87, Applicati	4.54e+00
34	20	7.9	56	4	PCT-US95-1	Sequence 100, Applicat	4.54e+00
35	20	7.9	74	4	PCT-US95-1	Sequence 99, Applicati	4.54e+00
36	20	7.9	81	4	PCT-US95-1	Sequence 98, Applicati	4.54e+00
37	20	7.9	423	1	US-08-470-	Sequence 174, Applicat	4.54e+00
38	20	7.9	669	3	US-08-896-	Sequence 3, Applicatio	4.54e+00
39	20	7.9	1140	3	US-08-896-	Sequence 1, Applicatio	4.54e+00
40	20	7.9	1305	3	US-08-896-	Sequence 37, Applicati	4.54e+00
41	19	7.5	66	4	PCT-US95-1	Sequence 93, Applicati	1.34e+01
42	19	7.5	260	3	US-08-594-	Sequence 45, Applicati	1.34e+01
43	19	7.5	1351	2	US-08-417-	Sequence 14, Applicati	1.34e+01
44	19	7.5	7218	2	US-08-232-	Sequence 13, Applicati	1.34e+01
45	19	7.5	9515	3	US-08-362-	Sequence 13, Applicati	1.34e+01

## ALIGNMENTS

1  
US-08-746-789A-1 STANDARD: DNA; UNC; 1920 BP.

AC xxxxxx

DE Sequence 1, Application US/08746789A

CC Sequence 1, Application US/08746789A

CC Patent No. 5789200

CC GENERAL INFORMATION:

CC APPLICANT: Ismail Kola, Martin J. Tyms, Christine Debouck

CC TITLE OF INVENTION: A No. 5789200e1 Human E1S Family Member, E1F3

CC NUMBER OF SEQUENCES: 4

CC CORRESPONDENCE ADDRESS:

CC ADDRESSEE: SmithKline Beecham Corporation

CC STREET: 709 Swedeland Road, P.O. Box 1539

CC CITY: King of Prussia

CC STATE: PA

CC COUNTRY: USA

CC ZIP: 19406-0939

CC COMPUTER READABLE FORM:

CC MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MB STORAGE

CC OPERATING SYSTEM: WINDOWS FOR WORKGROUPS

CC SOFTWARE: MICROSOFT WORD

CC CURRENT APPLICATION DATA:

CC APPLICATION NUMBER: US/08/746,789A

CC FILING DATE: No. 5789200e1 15, 1996

CC CLASSIFICATION: 514

CC PRIOR APPLICATION DATA:

CC APPLICATION NUMBER:

CC FILING DATE:

CC ATTORNEY/AGENT INFORMATION:

CC NAME: William T. Han

CC REGISTRATION NUMBER: 34,344

CC TELECOMMUNICATION INFORMATION:

CC TELEPHONE: 610 270 5219

CC TELEFAX: 610 270 4026

CC INFORMATION FOR SEQ ID NO: 1:

CC SEQUENCE CHARACTERISTICS:

CC LENGTH: 1920

CC TYPE: Nucleic Acid

CC STRANDEDNESS: Single

CC TOPOLOGY: Linear

CC ANTI-SENSE: No

FIGURE 1

CC	TYPE: nucleic acid
CC	STRANDEDNESS: unknown
CC	TOTAL COT:

STRANDEDNESS: unknown  
TOPOLOGY: unknown